

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/806, 634 A
Source: IFW16
Date Processed by STIC: 08/03/2005

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 08/03/2005

PATENT APPLICATION: US/10/806,634A

TIME: 15:10:00

Input Set : A:\SEQLIST_34909.TXT

Output Set: N:\CRF4\08032005\J806634A.raw

```

4 <110> APPLICANT: FRESNO ESCUDERO, Manuel
5   INIGUEZ PENA, Miguel Angel
7 <120> TITLE OF INVENTION: METHODS OF SCREENING FOR COX-2
8   INHIBITORS
10 <130> FILE REFERENCE: 34909-PCT-USA-I
12 <140> CURRENT APPLICATION NUMBER: US 10/806,634A
13 <141> CURRENT FILING DATE: 2004-03-23
15 <150> PRIOR APPLICATION NUMBER: 10/031047
16 <151> PRIOR FILING DATE: 2002-09-23
18 <150> PRIOR APPLICATION NUMBER: ES P9901557
19 <151> PRIOR FILING DATE: 1999-07-12
21 <160> NUMBER OF SEQ ID NOS: 6
23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 33
27 <212> TYPE: DNA
28 <213> ORGANISM: Artificial Sequence
30 <220> FEATURE:
31 <223> OTHER INFORMATION: Oligonucleotide primer
33 <400> SEQUENCE: 1
34 gggggatccg gattctaaca tggcttctaa ccc                               33
36 <210> SEQ ID NO: 2
37 <211> LENGTH: 30
38 <212> TYPE: DNA
39 <213> ORGANISM: Artificial Sequence
41 <220> FEATURE:
42 <223> OTHER INFORMATION: Oligonucleotide primer
44 <400> SEQUENCE: 2
45 gggagatctg gtaggctttg ctgtctgagg                                   30
47 <210> SEQ ID NO: 3
48 <211> LENGTH: 1979
49 <212> TYPE: DNA
50 <213> ORGANISM: Artificial Sequence
52 <220> FEATURE:
53 <223> OTHER INFORMATION: Human cyclooxygenase 2 promoter cloning fragment
W--> 55 <221> NAME/KEY: primer_bind
56 <222> LOCATION: (46)...(70)
57 <223> OTHER INFORMATION: Oligonucleotide No 1
W--> 59 <221> primer_bind
60 <222> LOCATION: (1925)...(1947)
61 <223> OTHER INFORMATION: Oligonucleotide No 1
W--> 63 <400> 3
64 gaattcagga ttgtaatgta aaatttttagt actctctcac agtatggatt ctaacatggc 60

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```

65 ttctaacc ca aactaacatt agtagctcta actataaaact tcaaattttca gtagatgcaa 120
66 cctactcctt taaaatgaaa cagaagattg aaattatttaa attatcaaaa agaaaatgat 180
67 ccacgctctt agttgaaatt tcatgtaaga ttccatgcaa taaataggag tgccataaat 240
68 ggaatgatga aatatgacta gaggaggaga aaggcttcct agatgagatg gaattttagt 300
69 catccgtgtc tcatgaagaa tcagatgtgt acactaagca aaacagttaa aaaaaaaacc 360
70 tccaagtgtg tctcttattt atttttttct tataagactt ctacaaattg aggtacctgg 420
71 tgtagtttta tttcaggttt tatgtgtgtca ttttcctgta atgctaagga cttaggacat 480
72 aactgaattt tctattttcc acttcttttc tgggtgtgtgt gtatatatat atgtatatat 540
73 acacacacac atatacatat atataatttt agtatctcac cctcacatgc tctccctga 600
74 gcactaccca tgatagatgt taaacaaaag caaagatgaa attccaactg ttaaaatctc 660
75 ccttccatct aattaattcc tcatccaact atgttccaaa acgagaatag aaaattagcc 720
76 ccaataagcc caggcaactg aaaagtaaat gctatgttgt actttgatcc atggtcacaa 780
77 ctcataatct tggaaaagtg gacagaaaag acaaaagagt gaactttaaa actcgaattt 840
78 attttaccag tatctcctat gaagggctag taaccaaatt aatccacgca tcagggagag 900
79 aaatgcctta aggcatacgt tttggacatt tagcgtccct gcaaattctg gccatcgccg 960
80 cttcctttgt ccatcagaag gcaggaaact ttatatgggt gacccgtgga gctcacatta 1020
81 actattttaca gggtaactgc ttaggaccag tattatgagg aggatttacc tttccgcct 1080
82 ctctttccaa gaaacaagga gggggtgaag gtacggagaa cagtatttct tctgttgaaa 1140
83 gcaacttagc tacaaagata aattacagct atgtacactg aaggtagcta tttcattcca 1200
84 caaaataaga gtttttttaa aaagctatgt tgtatgtcct gcatatagag cagatatata 1260
85 gcctattaag cgctgcactt aaacataaaa acatgtcagc ctttcttaac cttactcgcc 1320
86 ccagtctgtc ccgacgtgac ttctctgacc ctctaaagac gtacagacca gacacggcgg 1380
87 cggcggcggg agaggggatt cctgcgccc ccggacctca gggcgctca gattcctgga 1440
88 gaggaagcca agtgtccttc tgccctcccc cggatatcca tccaaggcga tcagtccaga 1500
89 actggctctc ggaagcgctc gggcaaagac tgcgaagaag aaaagacatc tggcggaaac 1560
90 ctgtgcgcct ggggcgggtg aactcgggga ggagaggag ggatcagaca ggagagtggg 1620
91 gactaccccc tctgtcccca aattggggca gcttctctggg tttccgattt tctcatttcc 1680
92 gtgggtaaaa aaccctgccc ccaccgggct tacgcaattt ttttaagggg agaggaggga 1740
93 aaaatttgtg gggggtacga aaaggcggaa agaaacagtc atttcgtcac atgggcttgg 1800
94 ttttcagtct tataaaaagg aaggttctct cggttagcga ccaattgtca tacgacttgc 1860
95 agtgagcgtc aggagcacgt ccaggaactc ctacgcagcg cctccttcag ctccacagcc 1920
96 agacgcccct agacagcaaa gcctaccccc cgcgcgcgcg cctgcccgcg gctgcgatg 1979
98 <210> SEQ ID NO: 4
99 <211> LENGTH: 70
100 <212> TYPE: DNA
101 <213> ORGANISM: Artificial Sequence
102 <220> FEATURE:
103 <223> OTHER INFORMATION: Cloning fragment where sequences of 5' and 3' ends
104 are shown in detail and the internal sequences are
105 represented by Ns.
W--> 108 <221> NAME/KEY: promoter
109 <222> LOCATION: (34)...(40)
110 <223> OTHER INFORMATION: Shorthand representation of sequence from
111 nucleotide 25 to nucleotide 882 of SEQ ID NO:6
W--> 113 <221> misc feature
114 <222> LOCATION: 34, 35, 36, 37, 38, 39, 40
115 <223> OTHER INFORMATION: n = A,T,C or G
W--> 117 <400> 4
W--> 118 ggggggatccg gattctaaca tggcttctaa cccnnnnnnn cctcagacag caaagcctac 60

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```

119 cagatctccc                                     70
121 <210> SEQ ID NO: 5
122 <211> LENGTH: 70
123 <212> TYPE: DNA
124 <213> ORGANISM: Artificial Sequence
126 <220> FEATURE:
127 <223> OTHER INFORMATION: reverse strand - Cloning fragment where sequences
128     of 5' and 3' ends are shown in detail and the
129     internal sequences are represented by Ns.
W--> 131 <221> NAME/KEY: promoter
132 <222> LOCATION: (34)...(40)
133 <223> OTHER INFORMATION: Shorthand representation of sequence from
134     nucleotide 25 to nucleotide 882 of SEQ ID NO:6
W--> 136 <221> misc_feature
137 <222> LOCATION: 31, 32, 33, 34, 35, 36, 37
138 <223> OTHER INFORMATION: n = A,T,C or G
W--> 140 <400> 5
W--> 141 gggagatctg gtaggctttg ctgtctgagg nnnnnnnnggg ttagaagcca tgtagaatc 60
142 cggatccccc                                     70
144 <210> SEQ ID NO: 6
145 <211> LENGTH: 1902
146 <212> TYPE: DNA
147 <213> ORGANISM: Homo sapiens
149 <400> SEQUENCE: 6
150 ggattctaac atggtcttcta acccaaaacta acattagtag ctctaactat aaacttcaaa 60
151 tttcagtaga tgcaacctac tcctttaaaa tgaaacagaa gattgaaatt attaaattat 120
152 caaaaagaaa atgatccacg ctcttagttg aaatttcatg taagattcca tgcaataaat 180
153 aggagtgcc aataatggaat gatgaaatat gactagagga ggagaaaggc ttcttagatg 240
154 agatggaatt ttagtcatcc gtgtctcatg aagaatcaga tgtgtacact aagcaaaaca 300
155 gttaaaaaaa aaacctccaa gtgagtctct tatttatttt tttcttataa gacttctaca 360
156 aattgaggta cctggtgtag ttttatttca ggttttatgc tgtcattttc ctgtaatgct 420
157 aaggacttag gacataactg aattttctat tttccacttc ttttctggtg tgtgtgtata 480
158 tatatatgta tatatacaca cacacatata catatatata ttttttagtat ctcacctca 540
159 catgctcttc cctgagcact acccatgata gatgttaaac aaaagcaaag atgaaattcc 600
160 aactgttaaa atctcccttc catctaatta attcctcatc caactatggt ccaaaacgag 660
161 aatagaaaat tagccccaat aagcccaggc aactgaaaag taaatgctat gttgtacttt 720
162 gatccatggt cacaactcat aatcttggaa aagtggacag aaaagacaaa agagtgaact 780
163 ttaaaaactc aatttatttt accagtatct cctatgaagg gctagtaacc aaaataatcc 840
164 acgcatcagg gagagaaatg ccttaaggca tacgttttgg acatttagcg tccctgcaaa 900
165 ttctggccat cgccgcttcc tttgtccatc agaaggcagg aaactttata ttggtgaccc 960
166 gtggagctca cattaactat ttacagggtg actgcttagg accagtatta tgaggaggat 1020
167 ttaccttttc cgctctctct tccaagaaac aaggaggggg tgaaggtagc gagaacagta 1080
168 tttcttctgt tgaaagcaac ttagctacaa agataaatta cagctatgta cactgaagg 1140
169 agctatttca ttccacaaaa taagagtttt ttaaaaagct atgtatgtat gtctgcata 1200
170 tagagcagat atacagccta ttaagcgctc tactaaaac ataaaacatg tcagcctttc 1260
171 ttaaccttac tcgccccagt ctgtcccgac gtgacttctc cgacctcta aagacgtaca 1320
172 gaccagacac ggcggcgggc ggcgggagag ggattccctg cgcggcgga cctcagggcc 1380
173 gctcagattc ctggagagga agccaagtgt cttctgccc tccccggta tcccatccaa 1440
174 ggcgatcagt ccagaactgg ctctcggaag cgctcgggca aagactgcga agaagaaaag 1500

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```
175 acatctggcg gaaacctgtg cgcctggggc ggtggaactc ggggaggaga gggagggatc 1560
176 agacaggaga gtggggacta cccctctgc tcccaaattg gggcagcttc ctgggtttcc 1620
177 gattttctca ttccgtggg taaaaaaccc tgccccacc gggcttacgc aattttttta 1680
178 aggggagagg agggaaaaat ttgtgggggg tacgaaaagg cggaaagaaa cagtcatttc 1740
179 gtcacatggg cttggttttc agtcttataa aaaggaaggt tctctcgggt agcgaccaat 1800
180 tgcatacga cttgcagtga gcgtcaggag cacgtccagg aactcctcag cagcgcctcc 1860
181 ttcagctcca cagccagacg ccctcagaca gcaaagccta cc 1902
```

RAW SEQUENCE LISTING ERROR SUMMARY
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Input Set : A:\SEQLIST_34909.TXT
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:4; N Pos. 34,35,36,37,38,39,40

Seq#:5; N Pos. 31,32,33,34,35,36,37

VERIFICATION SUMMARY

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Input Set : A:\SEQLIST_34909.TXT

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L:55 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:59 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:63 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:108 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:113 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:117 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:131 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:136 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5
L:140 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5
L:141 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0